Docket No.: PF-0420-2 DIV

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Narture

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Magna et al.

Title: HUMAN NUCLEOTIDE PYROPHOSPHOHYDROLASE-2

Serial No.: To Be Assigned Filing Date: Herewith

Examiner: To Be Assigned Group Art Unit: To Be Assigned

Official Draftsman

Commissioner for Patents Washington, D.C. 20231

SUBMISSION OF FORMAL DRAWINGS

Sir:

Transmitted herewith are Figure(s) 1A, 1B, 1C, 1D, 1E, 1F, 1G, 1H, 1I, 1J, 1K, 2A, 2B, 2C, 3A, and 3B, as sixteen (16) sheets of formal drawings for this application. Each sheet of drawing indicates the identifying indicia suggested in 37 CFR Section 1.84(c) on the reverse side of the drawings.

Applicants believe that no fee is due with this paper. However, if the Commissioner determines that a fee is necessary, the Commissioner is hereby authorized to charge any additional fees associated with this communication or credit any overpayment to Deposit Account No. **09-0108.** A duplicate copy of this communication is enclosed.

If there are any questions regarding the above, the Examiner is invited to call the undersigned at 650-855-0555.

Respectfully submitted,

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54 ATG M	108 GGG G	162 GAA E	216 GAG E	270 GAC D	324 CCG P	378 GTC V
၁၁၅	GCG A	CTG	GAA E	9 299	TGC	GCC
CCG	$ ext{CTG}$	9 299	TGG W	GAC D	GTG V	TCC
45 GCC	99 CAC H	153 CTG L	207 GAC D	261 GGC G	315 CGC R	369 CCG P
TCT	GCG	GCA A	GAG	GGA G	GCG	GCC CTG A L
CGC	GCT	ACT	CTG L	CCC	CCA	GCC A
36 GGA (90 GTC V	144 GCG A	198 GCC A	252 GAC CAC D H	306 TAC GGG Y G	360 TGG W
CTC	GTC V	ATG M	CCA P	GAC D	TAC Y	GAC D
TCC	TGT	CCA P	TCA	GTG V	TAC Y	ACG
27 ACC	81 CTC L	135 GAG E	189 CCC P	243 AAC N	297 TTC F	351 ACC T
550	TGT	GAG	CAG Q	TTC	CGC	CGC R
၁၁၅	CTC	ACC	9	TGG W	ATC	342 GAG GCG E A
18 GGA	72 CTG L	126 CCC P	180 ACC T	234 TCC S	288 GCC A	342 GAG E
၁၅၁	CCA	ACC	TAC Y	ACG	GCT	CTG
ე၅ე	CTG L	GCC	GTG V	TGG W	CTG	GCG
9 GCA (63 CTG L	117 GAC D	171 TCC S	225 GAG E	279 AGC S	333 CTG L
GCC CGA	TCG (CGA R	CGG R	AGC S	GAG	333 CGA CCG CTG C R P L A
သည	GCG	GCC	AGA R	GCC	TTC	CGA R
ب						

FIGURE 1A

432 GAG E	486 CTA L	540 GGG G	594 TGT C	648 TGC C	702 ACC T	756 GGC G
CGC R	CCA P	TGT	GCG A	999 v	GTC V	CCT
AAC N	TGC	AGC	GAT D	CCA P	GTG V	CAG Q
423 CTC L	477 CGC R	531 GGG G	585 GGG G	639 TGT C	693 TCG S	747 GAC D
TGC	TTC	TCG	GCT	CGG R	9	CGA R
TGG W	CGC	TGC	CCC	CCT	CTG	CTG
414 TTC F	468 GTG V	522 CCC P	576 AGC S	630 CGG R	684 CTC L	738 TCC S
9 292	CAC H	GGT	CCA P	GTG V	ATC	GTC V
CGC R	TAC Y	TGG W	TGC	TGC	CAC	AGG R
405 ACG T	459 AAC N	513 CCG P	567 CAC H	621 AAG K	675 GAC D	729 GCC A
CCC	TCC	385 385	CGC R	CAG Q	CCG	GGA
AAC N	TGC	TGG W	CGC R	GCG	TGC	CTA
396 TTG L	450 CGC R	504 GCG A	558 CGC R	612 GAG E	666 GAA E	720 CTG L
CAC H	CGC R	299	$ ext{TTG}$	CTG	TGT	CCA
GTG V	299	TGG W	CGC R	CCT	ACC	caa o
387 CGC R	441 CGT R	495 TCG S	549 CGT R	603 CGT R	657 GAC D	711 GGG G
GAG	A CCG P	GAA GCC E A	CCA GGC P G	5 d	CTT L	TCT
299	CAA	GAA E	CCA	CCC	AGC	CCA

FIGURE 1B

810 GCT A	864 GCC A	918 TTG L	972 CAG Q	.026 TCC S	1035 1044 1053 1062 1071 1080 TTC CAC AAT GGG ACC CTG CTG GAC AGG CGA GCT CAT GGG TAC GGG GCC CAC F H N G T L L D R R A H G Y G A H	1134 GCA A
TGT	GAG E	AAG K	9 299	1 TAC Y	GCC A	1 AAG K
GTC TGT V C	999 9	GAT D	GCT	AAA K	999	TGC
801 GGT G	855 GCA A	909 CTT L	963 GAG E	1017 AAG K	1071 TAC Y	1125 CAC H
CCT	TCT	ATC I	CGA R	CCC	999	TAC Y
GTG V	TTC F	ATC I	GTG V	ATG M	CAT H	ATC I
792 CGG R	846 GGC G	900 ACC T	954 CGA R	1008 CCC P	1062 GCT A	1116 GGC G
ACC TTC T F	GAT D	GTC V	TCC	ACC	CGA R	GCT
ACC	ATG M	GTG V	GAG E	999	AGG R	CAG Q
83 GA	837 CAG Q	891 TCT S	945 CCT P	999 TCC S	1053 GAC D	1107 GAC D
GCT CAC A H	828 C ATC AGG GCC C I R A Q	ATC I	CAC	GCC	CTG	CCA
GCT	AGG R	TCC	AAA K	AAA K	$_{\rm L}^{\rm CTG}$	CGC R
774 GAT D	828 ATC I	882 GGA G	936 GTG V	990 TGC C	1044 ACC T	1098 CTG L
AG. S	AAC N	AAC N	CTG	TGC		GGA
ACC T	GCC	GCC	TAC Y	$ ext{TTC}$	AAT N	CGG R
765 GCC A	819 CGC R	873 CAG Q	927 CCG P	981 ACT T	L035 CAC H	1089 CTS L
GTG V	AGC	G GCC	AAG K	GTG V	TTC	GAG E
ACT GTG T V	GAC	CAG Q	GAG	AAT N	TGG W	CTG

FIGURE 1C

188	.242	.296	1305 1314 1323 1332 1341 1350 CCC GAC ACC CGC TGC CTG GCA GGC TCC AGC CCC CGC TGC GGG GAC GCC P S L A G S S P R C G D A	.404	458	.512
GCC	CCT	TGT		CCT	CTG	CCG
A	P	C		P	L	P
1	1	1	1	1	1	1
CTT	CTC	CTC	GAC	TGC	TGT	GAG
L	L	L	D	C	C	E
GTA V	AAG K	9 299	999	CAC H	AAG K	999
1143 1152 1161 1170 1179 1188	1242 1233 1242	L287	1341	1395	1449	1503
TGG AAT GAG GCG GCG CTC ACT GTA CTT GCC	CCA GCC TGC GAC CCC CGA GAG TAC CTG ATC AAG CTC CCT	GTG	TGC	ATT	CAG	TCC
W N E A G A V R S G T A R L T V L A	P A C D P R P R E Y L I K L P	V	C	I	Q	S
CTC L	CTG L	J GAT D	cGC R	GAG E	TGC C	GAC D
CGG	TAC	CTG	CCC	AGG		GCT
R	Y	L		R	299	A
1170	1224	1278	1332	1386	1440	1494
GCC	GAG	TAC	AGC	AGA	TGT	GCT
A	E	Y	S	R	C	A
ACT T	1 CGA R	GCC A	TCC S	GAG E	GAG E	GTG V
9 299	CCC	CCT	999	CTG	GCA	GTT V
1161	1215	1269	1323	1377	1431	1485
TCG	CGG	GGC	GCA	CGT	GTG	CGT
S	R	G	A	R	V	R
CGC R	CCC P	AGT S	CTG L	CGC R	GTG V	ر 993 9
GTG	GAC	GGT	AGC	GTG	AAG	CGG
V	D	G		V	K	R
1152	1206	1260	1314	1368	1422	L476
GCC	TGC	CCA	CCC	TCT	GTG	GTC
A	C	P	P	S	V	V
GGT	GCC	CAG Q	TGC	TGC	CCA	CTG
GCG	CCA P	GGT	CGC R	TGC	CTC	999 9
1143	1197	1251	1305	1359	L413	L467
GAG	CAG	TGT	ACC	CGC	GTC	CGG
E	Q	C	T	R	V	R
AAT		GAC	GAC D	TCC	TAC Y	CCT
TGG W	1197 CCA GGC CAG C P G Q P	GAG	CCC	AGC	1990 1990	CCC

FIGURE 1D

566 TAC		1620 ; ACT T	.674 GAT D	1728 CCG P	L782 GAT D	1836 GAC D	1845 1854 1863 1872 1881 1890 GGC AAA CCC TAC TCG GGG CCT GTG GAG GCC CGG GTG ACG TTC GTG GAC CCC CGA G K P Y S G P V E A R V T F V D P R
GCC 1	A S	1 GTG V	1 TTT F	GCC A	GAA	GCC	CCC
ACC) H	GTG V	CCT	AAA K	$^{ m CTG}_{ m L}$	AGA R	GAC
557 TTC	A R I L G Q E P I G F T A Y	1584 1593 1602 1611 1620 TTT ACC ATT GAG GTG CCG CCC TCC ACC CAG CGG CTG GTG GTG ACT F T I E V P P S T Q R L V V T	1629 1638 1647 1656 1665 1674 TTT GTG GAC CCC AGC GGT GAG TTC ATG GAC GCT GTC CGG GTC TTG CCT TTT GAT F V D P S G E F M D A V R V L P F D	1683 1692 1701 1710 1719 1728 CCT CGA GGT GCC GGC GTG TAC CAC GAG GTC AAG GCC ATG CGG AAG AAA GCC CCG PR G A G C CCG A G C C CCG A G C C C C C C	1737 1746 1755 1764 1773 1782 GTC ATT TTA CAT ACC AGC CAG AGC AAC ACG ATC CCC CTG GGC GAG CTG GAA GAT V I L H T S Q S N T I P L G E L E D	1791 1800 1809 1818 1827 1836 GAG GCG CCC CTG GGC GTC CTG CCT TCT GGC GCT TTC CGC AGA GCC GAC E A P L G E L V L P S G A F R A D	1881 GTG V
1) 0 0	1 CGG R	1 GTC V	CGG R	ر 1990 19	TTC	TTC
ATC	H	CAG Q	CGG R	ATG M	CTG L	GCT A	ACG
548) G	.602 ACC T	.656 GTC V	.710 GCC A	1764 CCC P	1818 GGC G	1872 GTG V
ן טעט	E E	1 TCC S	1 GCT A	1 AAG K	ATC I	TCT S	CGG.
ر د د	CAG Q	CCC	GAC D	GTC V	ACG	CCT	GCC
539	ر رود رود	.593 CCG	.647 ATG M	.701 GAG E	L755 AAC N	1809 CTG L	1863 GAG E
1	L	1 GTG V	1 TTC F	1 CAC H	AGC S	GTC V	GTG
Č E	i CI	GAG	GAG E	TAC	CAG Q	${ m CTG}$	CCT
530	ATT	.584 ATT I	.638 GGT G	.692 GTG V	1746 AGC S	L800 GAG E	1854 GGG G
1	AGG R	1 ACC T	1 AGC S	1 GGC G	ACC T	299 9	TCG
7	P CCC	TTT F	CCC	GCC	CAT H	CTG	TAC
521	TTC	1575 GAC D	.629 GAC D	.683 GGT G	1737 TTA L	1791 CCC P	1845 CCC P
, ,	CGC R	157 GGC GA G D	1 GTG V	1 CGA R) ATT I	GCG A	AAA K
!	CTA CGC TTC C	CAG Q	TTT F	CCT	GTC	GAG	9 299

FIGURE 1E

44 AC	98 CG	52 CC	06 AC	90 09	14 TC	68 TG
19, G,	199. A	20 G A	21 A 2	21 2 T S	22 3 A H	22 2 G
AGC S	CGT R	GCC A	CTC	TCC	GA(E	TT(F
GAC D	CTC	GTG V	TCG S	999	GTG V	TGC
935 GTG V	989 GAC D	043 CGG R	.097 TGG W	151 GAG E	AAC N	:259 CGC R
1917 1926 1935 1944 TCT GCC CCC AGT GAC CTG CGC TTC GTG GAC AGC GAC S A P S D L R F V D S D	1962 1971 1980 1989 1998 GCT CCA CTG CGC AC ATG TTC TCC GTG GAC CTC CGT GCG A P L R T Y G M F S V D L R A	2025 2034 2043 2052 CTG CAG GTG GCG GTG CGG GTG GCC GCC L Q V G P V A V R V A A	2070 2079 2088 2097 2106 CAC ATG CCA GGC CAC GTG GAG GCC CTC AAG CTG TGG TCG CTG AAC H M P G H V E A L K L W S L N	2124 2151 2160 GGC TTG TGG GAG GAG AGC GGC TTC CGG CGC GAG GGG TCC TCG G L W E E E S G F R R E G S S	2196 2205 2214 TTC CTG GTG GGC AAC GTG GAG ATC F L V G N V E I	2232 2241 2250 2259 2268 CGC CTG TTC AAT CTG GAC GTG CCT GAG CGC CGC TGC TTC GTG R L F N L D V P E R R C F V
CGC R	TCC	GCC	AAG K	CGG R	GTG V	CGC R
926 CTG L	980 TTC F	.034 GTG V	:088 CTC L	1142 TTC F	1196 CTG L	2250 GAG E
1 GAC D	1 ATG M	CCG P	GCC A	2 GGC G	TTC F	CCT
AGT S	9 299	999	GAG	AGC	GTC V	GTG V
917 CCC P	971 TAC Y	.025 GTG V	.079 GTG V	:133 GAG E	2178 2187 GTG CGC CGG GAG GAG CGC GTC V R R E E R V	2241 GAC D
1 GCC A	1 ACC T	2 CAG Q	2 CAC H	2 GAG E	GAG E	CTG
TCT	CGC	$^{ m CTG}$	9 299	GAG E	GAG	AAT N
1908 GCG A	.962 CTG L	2016 CAG Q	2070 CCA P	:124 TGG W	2178 CGG R	2232 TTC F
1908 TCG GCG GCG T S A A S	1 CCA P	2016 GCG GAG CAG C A E Q I	2 ATG M	TTG	CGC R	CTG L
TCG	GCT	GCG A	CAC H	9	GTG V	CGC R
1899 ACC T	1953 . CTG L	2007 TCC S	061 ATC I	2115 ACC T	2169 CGG R	2223 CGG R
1 CTC L	1 GAG E	2 86C 6	CAG Q	GAG E	2169 GGC CCC CGG G P R	2223 3 GAG CGG C E R F
1899 GAC CTC ACC ' D L T	1953 GGC GAG CTG C G E L A	2007 CCC GGC TCC G P G S A	2061 AGC CAG ATC CA S Q I H	2115 CCC GAG ACC G P E T G	9 299	CGG R

FIGURE 1F

- (3.6)	ω υ	o U	C 4	œ ڻ ع	7 D	9 9
2322 GGC G	2376 CCC P	2430 CTC L	2484 GCC A	253 CC P	259 AC T	2646 AAG K
GAG	2 AAC N	TGC C	ACC	CTC	CGG R	GCC A
GTG V	GCC	GCC	2484 GTC ACC GCC V T A	CCA P	CGT R	CTC
2304 2313 2322 TTC ACC CCC AGC GAG GTG GAG GGC F T P S E Q V E G	2358 2367 CCC GCC CCC GGC TTC TCC P A P G F S	2394 2403 2412 2421 2430 CGC TTT GAC AGC GCG GTC ACC GGC CCC AAT GGC GCC TGC CTC R F D S A V T G P N G A C L	2457 2466 2475 GAC AGG CCA GAC GCC TAC ACC GCC CTG D R P D A Y T A L	2502 2511 2520 2529 2538 GAG GAG CCG GCC CCT TCC TTG CCC CGC CCA CTC CCG E E L E P A P S L P R P L P	2565 2574 2583 2592 CAG CCC TAC CTG GAC AGG CTG GGG TAC CGT CGG ACG Q P Y L D R L G Y R T	2619 2628 2637 2646 TTC AAG CGT AAC GGC TTC CGC ATC AAC CTC GCC AAG F K R N G F R I N L A K
2 GAG E	TTC F	AAT N	GCC	CCC		ATC
AGC S	9 299	CCC	ACC	${ m TTG}$	CTG L	CGC R
2304 CCC P	2358 : CCC P	2412 GGC G	2466 TAC Y	1520 TCC S	2574 AGG R	2628 TTC F
2 ACC T	gcc A	ACC T	GCC A	CCT P	GAC D	
TTC	CCC	GTC V	GAC D	GCC	CTG	AAC N
2295 AAC GAC AAG 1 N D K F	2349 AAT CTG GAG (N L E I	403 GCG A	2457 ; CCA P	511 CCG P	2565 TAC Y	2619 ; CGT R
2 GAC D	2 CTG L	2 AGC S	2 AGG R	GAG E	CCC	, AAG K
AAC N	AAT N	GAC D	GAC D	CTG	CAG Q	TTC
2286 : GCC A	2340 . GTC V	394 TTT F	2448 GCC	:502 GAG E	2556 ACC T	2610 GCC A
TAC Y	CTG	2 CGC R	GAC		2556 GTC ACC V T	2610 CCC GCC 7 P A I
GCC	ACG	3 299	TGC	9 9	၁၅၁	GAT D
	2331 GTC V	2385 TGG W		2493 GGC G	2547 GTG	2601 GAC D
2277 AAG GTG CGC K V R	2331 GTG GTG GTC V V V	2385 CGT GCC TGG R A W	GCC A	2493 ACC CTG GGC T L G	2547 GCC ACC GTG A T V	2601 GAC CAC GAC D H D
AAG K	GTG V	CGT R	2439 CCC GCC TTC P A F	ACC T	GCC	GAC

FIGURE 1G

2700 TGG CGC AGC W R S ,	2754 TTC GCC AGG F A R	2781 2790 2799 2808 GAG TAC AAC GTG GTC CCC TTC CGA GAG GGC ACA CCT E Y N V V P F R E G T P	2853 2862 AAC CCG CAG GAG TTC CGG N P Q E F R	2889 2898 2907 2916 AAG ATC CAG GGT CCC CAG GAG TAT ATG GTC CGC TCC K I Q G P Q E Y M V R S	2961 2970 CAG CTC TAC GGA CTT CGG Q L Y G L R	2988 2997 3006 3015 3024 AGT GTG CGA GCC CCC GAG CGT CCG GGC ACC TCG GCA GCC TGC GTG S V R D P E R P G T S A A C V
TGG W	TTC	9 9	GAG E	GTC V	GGA	GCC
2691 TAC CCG Y	2745 ACT GCC AGC CAC TTC CGC T A S H F R	2799 . GAG E	2853 CAG Q	2907 ATG M	2961 TAC Y	3015 GCA A
2 TAC Y	TTC	CGA R	CCG	TAT	CTC	TCG S
3TG J	CAC H	${ m TTC}$	AAC	GAG E	CAG Q	ACC T
2682 GGG CCT (G P	736 AGC S	2790 CCC P	2844 CCC P	2898 CAG Q	2952 GGC G	3006 GGC G
2 GGG G	GCC A	GTC V	TGG W	CCC	CGC R	GCG.
LAT I	ACT T	GTG V	TGG W	GGT G	ACC	CGT R
2673 GCC GAG GCC <i>P</i> A E A D	2718 2727 CAG GGG GCC CCG GTG Q G A P V	2781 TAC AAC Y N	2835 2844 CTC CTG GCC TGG TGG CCC L L A W W P	2889 ATC CAG I Q	2934 2943 2952 GGC AGC CAC CCA CGC ACC G S H P R T R G	2997 CCC GAG P E
GCC O	GCC	GAG	CTC	AAG K	CAC H	GAC
664 CCC P	2718 . GGG G	2772 AAG TAC K Y	2826 GGC GAT G	2880 AAG GTG K	2934 AGC S	2988 CGA R
2 GAC D			2 66C 6	AAG K	, 299 9	GTG V
GGT G	TGC	GAC	ACT T	TC	999	
2655 CCA P	2709 GAA E	2763 GCG A	2817 TGG	2871 TTC F	2925 AAC GCA GGG N A G	2979 CGG R
2 AGG R	2709 CTG CGG GAA L R E	GAG	TCC	TGC	AAC	2979 GAT GCC CGG D A R
2655 CCC AGG CCA P R P	CTG	GTG	GCC	2871 GCC TGC TTC C A C F I	CAC	GAT

FIGURE 1H

3078 CTG	1132 CTT L	3186 GCC A	3240 ACT T	3294 TTT F	3348 GCC A	3402 AGG R
3 ACG T	CTC	GCT A	TAC	TGC	ACA	CAG
AGG R	GGA	CCA	GTC V	CGC R	9 9	TTC F
3069 3078 CAG GTG GAC AGG ACG CTG Q V D R T L	3105 3114 3123 3132 ATG CCC CAG GGC AGC TGC CGG CGC GTG GCC GTC AAC GGA CTC CTT M P Q G S C R R V A V N G L L	3150 3159 3168 3177 3186 CTG ACC CGC CCA CCG GTG CCC GCG GAG GAC CCA GCT L T R H P P V P A E D P A A	3195 3204 3213 3222 3231 3240 TCC ATG CTC CTC CTC CTC CTC CTC CTC CTC CTC C	3249 3258 3267 3276 3285 3294 ACT GAC CAG AGC CTTG GCC AAG GAG ATC GCC ATT GGC CGC TGC TTT T D Q S P R L A K E I A I G R C F	3303 3312 3321 3330 3339 3348 GGT TCC TCT GAC GGY TTC TCC AGA GAG ATG AAG GCT GAT GCC GGC ACA GCC G S S D G F S R E M K A D A G T A	3357 3366 3375 3384 3393 3402 GTC ACC TTC CAG TGC CGG GAG CCG GCC GGA CGA CCC AGC CTC TTC CAG AGG V T F Q C R E P P A G R P S L F Q R
GTG V	GTC V	GAG E	TAT Y	ATT	GAT	AGC S
CAG Q	GCC	GCG A	AAC N	GCC	GCT A	CCC
3042 3051 3060 TGC AGC GGG ATG CTG TTC GAC CAG CGG C S G M L F D Q R	3114 GTG V	3168 CCC P	3222 CAC H	3276 ATC I	3330 AAG K	3384 CGA R
3 CAG Q	CGC R	GTG V	355 9	GAG	ATG	GGA
GAC D	CGG R	CCG	CTG	AAG K	GAG E	GCC
1051 TTC F	1105 TGC C	3159 CCA P	3213 CCT P	3267 GCC A	3321 AGA R	3375 CCG P
3 CTG L	3 AGC S	S CCC P	GAC D	TTG L	TCC	CCA
ATG M	9 299	CAC H	\mathtt{CTA}	CGC R	TTC	GAG
042 GGG G	096 CAG Q	150 CGG R	204 CCC P	258 CCA P	3312 GGY G	3366 CGG R
3 AGC S	3 CCC P	3 ACC T	3 GCC A	3 AGC S	GAC D	TGC
TGC	ATG M	CTG L	$ ext{CTG}$	CAG Q	TCT	CAG Q
3033 AAG K	3087 ATT I	141 TAC Y	195 ATG M	(249 GAC D	303 TCC S	3357 TTC F
TTC	ACC	3141 3 GAT TAC C D Y L	TCC	3 ACT T	GGT G	ACC
GAG	GTG V	CGG R	TTC F	GTC	GAT D	GTC

FIGURE 11

3456 AGC GAG S E	3501 3510 CGC ACC CGC GGT AGG R T R R G R	3564 CAG ACT CCT	3618 CCC AGG	3663 3672 GTC AGA CAA GAA CCC	3726 TAT GGA	3771 3780 CGA CGT TCA ACC CTA	3834 CGT GCG TAT
ATG M	CGG R	CAG	CTC	CAA	TGG	TCA	
3447 CGC AGG GAG R R E	3501 ACC CGC T R	3555 TCC CTC	3609 AGC CCC	3663 AGA	3717 GCT GTG	3771 CGT	3825 ACT TCT
AGG R	ACC T	TCC	AGC		GCT	CGA	
CGC R		ACC	GAC	AGA	${ m GTT}$	GGA	CTG
3438 : ATC I	3492 CCC CTC P L	3546 TTT CCC ACC	3600 TCT CCA GAC	3654 GAA CTC	3708 ATT	3762 CCC AGG	3816 CCG TCT
3438 GAC ATC D I					ACA		
GGT	GGT G	၁၅၁	TCT	CCA	AAG	GCA	AAG
3429 GCA CTT A L	3483 CGG GCC TCA GGT R A S G	3537 3GG CCT	3591 CCT CCT	3645 CCC TTT	3699 3708 ACC AGG AAG ACA ATT GTT	3753 GGG CCA GCA	3807 GCT CAG
ACA (T	CGG (CAG () သည		AAC 7	CTG (CCA
3420 3429 3438 TCC CCG GCG ACA GCA CTT GGT GAC ATC S P A T A L G D I	3474 GCA CAG GCC A Q A	3528 3537 CCT GGG CAG GGG CCT CGC	3582 AGT TTT	3636 CTT TCC CGC	3690 GGT AGA AAC	3744 GTG ACT	3798 CCG CTC
TCC	GCA A	TGA	GGA	၁၃၃	GAT	909	GAC
3411 CTG CTG GAG L L E	3465 GCG GCG CAG A A Q	3519 GTC CGG CAG V R Q	3573 TTG ACC CCA	3627 TGT CTG GGT	3681 AGA GCA TCC	3735 GAG TTT	3789 TGA AGG
CTG	GCG A	GTC	TTG	TGT	AGA	ATG	၁၁၅

FIGURE 1J

3888 ACA AAG	3942 GAG CTT	3996 GTG TGA	4050 TCC CAT	4104 CTT AGA	4158 GCT TTA
299	သသ	TGG	\mathtt{GGT}	GTG	AAT
3879 GCG TTT	3933 CCC TGC	3987 TGG TTG	4041 GGA AGG	4095 TGT TTG	4149 ATT AAA AAT
CTG	GTC	AGC	GGA	ACG	TGA
3870 GGA GTT	3924 GAC CAG GTC	3978 GTG GGG	4032 TGA AAA	4086 ATT CTC	4140 GGG GGC
	CTA	AGT	CAG Q	ACA	TCC
3861 CTA CCC TTG	3915 TCA GTG	3969 AAA CAA AGT	4014 4023 GGT CCA GGC TGG GCC CAG G P G W A Q	4077 AGG GGA ACA	4131 AGG CCC
CTT	AGC	TTG	9 299	CTG	299
3852 TTC AAT	3906 GTT TGG	3960 TAT TTA TTG	4014 GGT CCA G P	4068 CTG GGG	4122 GCG TTG
GAT	CCT	GGT	TGG	GCT	999
3843 TTT GAC CCT	3897 TCC CCT CTG	3951 TGT TTT TGG	4005 GTG GGG GTG V G V	4059 GCG GGG GAG	4113 GAC CTG CCC GGG
TTT	TCC	TGT	GTG V	909	GAC

FIGURE 1K

TTT CCA AAA AAA AAA AAA AAA NAA A 3'

4176

	MAS LLPLLCLCVVAAHLAGARDATPTEEPMATALGLNIPH-2 MVGTKAWVFSFLVL = EVTSVLG RQTMLTQSVRRVQPGK NIPH-1	I-2 I-1
2 8	ERRSVYTGOPSPALEDWEEASEWTSWFNVDHPGGDGDFES NIPPH-2 KNPSIFA - KPADTLE SPGEWTTWFNIDYPGGKGDYER NIPPH-1	1-2 1-1
7 4	LAAIRFYYGPARVCPRPLALEARTTDWALPSAVGERVHLN NTPH-2 LDAIRFYYG-DRVCARPLRLEARTTDWTPAGSTGQVVHGS NTPH-1	1-2 1-1
17	PTRGFWCLNREQPRGRRCSNYHVRFRCP LEASW NTPPH-2 PREGFWCLNREORPGONCSNYTVRFLCPPGSLRRDTERIW NTPPH-1	1-1
52	GAWGPWGPCSGSCGP-GRRLRRHCPSPAGDACPGRPLEANTPH-2 SPWSPWSKCSAACGQTGVQTRTRICLAEMVSLCSEASEEG NTPH-1	1-7
88 89	QKCVRPRCPGCSL DTCECPDHILLGSVVT NIPPH-2 QHGMGQDCTACDLTCPMGQVNADCDACMCQDFMLHGAVSL NIPPH-1	+ +
118 33	PSGOPLLGARVSLRDQ-PGTVATSDAHGTFRVPGVCADSR MTPFH-2 PGGAPASGAAIYLLTKTPKLLTQTDSDGRFRIPGLCPDGK NTPFH-3	
257 273	ANIRAOMDGFSA - GEAOAOANGSISVVTIILDKLEKPYLV NTPH-2 SILKITKVKFAPIVLTMPKTSLKAATIKAEFVRAETPYMV NTPH-3	
396	KHPESRVREAGONVTFCCKASGTPMPKKYSWFHNGTLLDRNIPPH-2 MNPETKARRAGOSVSLCCKATGKPRPDKYFWYHNDTLLDP NIPPH-	H H
336	RAHGYGAHLELRGLRPDQAGIYHCKAWNEAGAVRSGTARL NIPPH-S SLYKHESKLVLRKLQQHQAGEYFCKAQSDAGAVKSKVAQL NIPPH-	

FIGURE 2A

FIGURE 2B

376 TVLAPGOPACDPRPREYLIKLPEDCGOPGSGPAYLDVGLC NIPPH-2	456 PVKVVAECGCOKCLPPRGLVRGRVVAADSGEPLRFARILL MTPPH-	496 GQEPIGFTAYQGDFTIEVPPSTORLVVTFVDPSGEFMDAV NTPRH-	536 RVLPFDPRGAGVYHEVKAMRKKAPVILHTSOSNTIPLGEL NIPPH-	576 EDEAPLGELVLPSGAFRRADGKPYSGPVEARVTFVDPRDLINTPRH-	616 TSAASAPSDLRFVDSDGELAPLRTYGMFSVDLRAPGSAEQ NTPPH-	656 LOVGPVAVRVAASQIHMPGHVEALKLWSLNPETGLWEEES NTPFH-	696 GFRREGSSGPRVRREERVFLVGNVEIRERRLFNLDVPERR	736 RCFVKVRAYANDKFTPSEQVEGVVVTLVNLEPAPGFSANP NTPH-2
393 IVIASDETPCNPVPESYLIRLPHDCFQNATNSFYYDVGRC NIPPH-1	473 PTKVAKECSCORCTETRSIVRGRVSAADNGEPMRFGHVYM MTPPH-	513 GNSRVSMTGYKGTFTLHVPQDTERLVLTFVDRLQKFVNTT NTPRH-	553 KVLPFNKKGSAVFHEIKMLCRKEPITLEAMETNIIPLGEV NIPPH-	593 VGEDPMAELEIPSRSFYRQNGEPYIGKWKASVTFLDPRNINTPRH-	633 STATAAQTDLNFINDEGDTFPLRTYGMFSVDFRDEVTSEP NTPPH-	673 LNAGRVRVHLDSTQVKMPEHISTVKLWSLNPDTGLWEEEG NTPFH-	713 DFKFENQRRNKREDRTFLVGNLEIRERRLFNLDVPESR NTPFH	751 RCFVKVRAYRSERFLPSEQIQGVVISVINLEPRTGFLSNP NTPH-1
		PVKVVAECGCOKCLPPRGLVRGRVVAADSGEPLRFARI PTKVAKECSCORCTETRSIVRGRVSAADNGEPMRFGHV	PVKVVAECGCQKCLPPRGLVRGRVVAADSGEPLRFARILL PTKVAKECSCORCTETRSIVRGRVSAADNGEPMRFGHVYM GQEPIGFTAYQGDFTIEVPPSTORLVVTFVDPSGEFMDAV GNSRVSMTGYKGTFTLHVPQDTERLVLTFVDRLQKFVNTT	PVKVVARECSCORCTETRSIVRGRVVAADSGEPLRFARILL GOEPIGFTAYOGDFTIEVPPSTORLVVTFVDPSGEFMDAV GNSRVSMTGYKGTFTLHVPQDTERLVLTFVDRLQKFVNTT RVLPFDPRGAGVYHEVKAMRKKAPVILHTSOSNTIPLGEL KVLPFNKKGSAVFHEIKMLCRKEPITLEAMETNILPLGEV	PVKVVABECGCOKCLPPRGLVRGRVVAADSGEPLRFARILLI PTKVARKECSCORCTETRSIVRGRVSAADNGEPMRFGHVYM GOEPIGFTAYOGDFTIEVPPSTORLVVTFVDPSGEFMDAV GNSRVSMTGYKGTFTLHVPPQDTERLVLTFVDRLQKFVNTT RVLPFDPRGAGVYHEVKAMRKKAPVILHTSOSNTIPLGEL KVLPFNKKGSAVFHEIKMLCRKEPITLEAMETNILPLGEV EDEAPLGELVLPSGAFRRADGKPYSGPVEARVTFVDPRDL VGEDPMAELFIPSRSFYRQNGEPYIGKVKASVTFLDPRNI	PVKVVAECGCORCTETRSIVRGRVVAADSGEPLRFARILLI PTKYARECSCORCTETRSIVRGRVSAADNGEPMRFGHVYM GOEPIGFTAYOGDFTIEVPPSTORLVVTFVDPSGEFMDAV GNSRVSMTGYRGTFTLHVPQDTERLVLTFVDPRLORLOREL RVLPFDPRGAGVYHEVRAMRKKAPVILHTSOSNTIPLGEL KVLPFDRKGSAVYHETRRADGKPYSGPVEARVTFVDPRDL CDEAPLGELVLPSGAFRRADGKPYSGPVEARVTFVDPRDL VGEDPMAEIFIPSRSFYRQNGEPYIGKWKASVTFVDPRDL TSAASAPSDLRFVDSDGELAPLRTYGMFSVDFRDEVTSEP	PVKVVA RECSCORCTETRSIVRGRVSAADNGEPMRFGHVYM PTKVARECSCORCTETRSIVRGRVSAADNGEPMRFGHVYM GOEPIGFTAYOGDFTIEVPPSTORLVVTFVDPSGEFMDAV GNSRVSMTGYRGTFTLHVPQDTERLVLTFVDRLOKFVNTT RVLPFNRKGSAVFHEIKMLCRKEPITLEAMETMILPLGEL KVLPFNRKGSAVFHEIKMLCRKEPITLEAMETMILPLGEL TSAASAPSDLRFVDSDGELAPLRTYGMFSVDFRDENI TSAASAPSDLRFVDSDGELAPLRTYGMFSVDFRDEVTSEP STATAAQTDLNFINDEGDTFPLRTYGMFSVDFRDEVTSEP LOVGPVAVRVAASQIHMPGHVEALKLWSLNPFTGLWEEES LOVGPVAVRVAASQIHMPGHVEALKLWSLNPFTGLWEEES	PVKVVABECGCORCTEPRGLVRGRVVAADSGEPLRFARILL PTKVAARECSCORCTETRSIVRGRVVAADSGEPLRFARILL GOEPIGFTAYOGDFTIEVPPSTORLVVTFVDPSGEFMDAV GNSRVSMTGYKGTFTLHVPQDTERLVLTFVDPSGEFMDAV GNSRVSMTGYKGTFTLHVPQDTERLVLTFVDRLOKEVNTT RVLPFNRKGSAVFHEIKMLCRKEPITLEVPSGEFMDAV TSAASAPSDLRFYDSDGELAPLRTYGRETNILLEAMETNILPLGEV TSAASAPSDLRFVDSDGELAPLRTYGRFSVDLRAEGSAEQ STATAAAQTDLNFINDEGDTFPLRTYGMFSVDLRAEGSAEQ LOVGPVAVRVAASQIHMPGHVEALKLWSLNPPTGLWEEES LOVGPVAVRVHLDSTOVKMPEHISTVKLWSLNPDTGLWEEES GFRREGSSGPRVRREERVFLVGNVEIRERRLDVDVPERR

FIGURE 2C



